1600

RAW SEQUENCE LISTING   DATE: 01/15/2002
PATENT APPLICATION: US/09/597,796B   TIME: 12:32:10

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Output Set: N:\CRF3\01152002\I597796B.raw

Does Not Comply
Corrected Diskette Needed

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 3 <110> APPLICANT: Skeiky, Yasir
 4         Reed, Steven
 5         Alderson, Mark
 6         Corixa Corporation
 8 <120> TITLE OF INVENTION: Fusion Proteins of Mycobacterium Tuberculosis
10 <130> FILE REFERENCE: 014058-009050US
12 <140> CURRENT APPLICATION NUMBER: US 09/597,796B
13 <141> CURRENT FILING DATE: 2001-06-20
15 <150> PRIOR APPLICATION NUMBER: US 09/056,556
16 <151> PRIOR FILING DATE: 1998-04-07
18 <150> PRIOR APPLICATION NUMBER: US 09/223,040
19 <151> PRIOR FILING DATE: 1998-12-30
21 <150> PRIOR APPLICATION NUMBER: WO PCT/US99/07717
22 <151> PRIOR FILING DATE: 1999-04-07
24 <150> PRIOR APPLICATION NUMBER: US 09/287,849
25 <151> PRIOR FILING DATE: 1999-04-07
27 <150> PRIOR APPLICATION NUMBER: US 60/158,338
28 <151> PRIOR FILING DATE: 1999-10-07
30 <150> PRIOR APPLICATION NUMBER: US 60/158,425
31 <151> PRIOR FILING DATE: 1999-10-07
33 <160> NUMBER OF SEQ ID NOS: 30
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38 <211> LENGTH: 588
39 <212> TYPE: DNA
40 <213> ORGANISM: Mycobacterium tuberculosis
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43 <223> OTHER INFORMATION: Ra35, N-terminus of MTB32A (TbRa35FL)
45 <220> FEATURE:
46 <221> NAME/KEY: CDS
47 <222> LOCATION: (1)..(588)
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53 aacaacgccg tgggcgccgg gaccggcatc gtcatcgatc ccaacggtgt cgtgctgacc 180
54 aacaaccacg tgatcgcggg cgccaccgac atcaatgcgt tcagcgtcgg ctccggccaa 240
55 acctacggcg tcgatgtggt cgggtatgac cgcacccagg atgtcgcggt gctgcagctg 300
56 cgcggtgccg gtggcctacc atcggcggcg atcggtggcg gcgtcgcggt tggtgagccc 360
57 gtcgtcgcga tgggcaacag cggtgggcag ggcggaacgc ccgtgcggt gcctggcagg 420
58 gtggtcgcgc tcggccaaac cgtgcaggcg tcggattcgc tgaccggtgc cgaagagaca 480
59 ttgaacgggt tgatccagtt cgatgccgcg atccagcccg gtgattcggg cgggcccgtc 540
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64 <211> LENGTH: 195
65 <212> TYPE: PRT
66 <213> ORGANISM: Mycobacterium tuberculosis
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67 <223> OTHER INFORMATION: Ra35, N-terminus of MTB32A (TbRa35FL)
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75 Asn Ile Asn Thr Lys Leu Gly Tyr Asn Asn Ala Val Gly Ala Gly Thr
76   35   40   45
77 Gly Ile Val Ile Asp Pro Asn Gly Val Val Leu Thr Asn Asn His Val
78   50   55   60
79 Ile Ala Gly Ala Thr Asp Ile Asn Ala Phe Ser Val Gly Ser Gly Gln
80 65   70   75   80
81 Thr Tyr Gly Val Asp Val Val Gly Tyr Asp Arg Thr Gln Asp Val Ala
82   85   90   95
83 Val Leu Gln Leu Arg Gly Ala Gly Gly Leu Pro Ser Ala Ala Ile Gly
84   100   105   110
85 Gly Gly Val Ala Val Gly Glu Pro Val Val Ala Met Gly Asn Ser Gly
86   115   120   125
87 Gly Gln Gly Gly Thr Pro Arg Ala Val Pro Gly Arg Val Val Ala Leu
88   130   135   140
89 Gly Gln Thr Val Gln Ala Ser Asp Ser Leu Thr Gly Ala Glu Glu Thr
90 145   150   155   160
91 Leu Asn Gly Leu Ile Gln Phe Asp Ala Ala Ile Gln Pro Gly Asp Ser
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96   195
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100 <211> LENGTH: 1872
101 <212> TYPE: DNA
102 <213> ORGANISM: Mycobacterium tuberculosis
104 <220> FEATURE:
105 <223> OTHER INFORMATION: MTB32A (TbRa35FL) cDNA
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110 gtcatggttg ctgagcgtgc tggctgccgt cgggctgggc ctggccacgg cgccggccca 180
111 ggcggccccg ccggccttgt cgcaggaccg gttcgccgac ttccccgcgc tgcccctcga 240
112 cccgtccgcg atggtcgccc aagtggcgcc acaggtggtc aacatcaaca ccaaactggg 300
113 ctacaacaac gccgtgggcg ccgggaccgg catcgtcatc gatcccaacg gtgtcgtgct 360
114 gaccaacaac cacgtgatcg cgggcgccac cgacatcaat gcgttcagcg tcggctccgg 420
115 ccaaacctac ggcgtcgatg tggtcgggta tgaccgcacc caggatgtcg cggtgctgca 480
116 gctgcgcggt gccggtggcc tgccgtcggc ggcgatcggt ggcggcgtcg cggttggtga 540
117 gcccgtcgtc gcgatgggca acagcggtgg gcagggcgga acgccccgtg cggtgcctgg 600
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119 gacattgaac gggttgatcc agttcgatgc cgcaatccag cccggtgatt cgggcgggcc 720
120 cgtcgtcaac ggcctaggac aggtggtcgg tatgaacacg gccgcgtccg ataacttcca 780
121 gctgtcccag ggtgggcagg gattcgccat tccgatcggg caggcgatgg cgatcgcggg 840

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       137 tacggtgact attacgtgtg gagcgacacc agcgagcgct acaccgacgc ccggatcatc 1800
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       154              20                  25                  30
       155 Ala Pro Pro Ala Leu Ser Gln Asp Arg Phe Ala Asp Phe Pro Ala Leu
       156          35                  40                  45
       157 Pro Leu Asp Pro Ser Ala Met Val Ala Gln Val Ala Pro Gln Val Val
       158      50                  55                  60
       159 Asn Ile Asn Thr Lys Leu Gly Tyr Asn Asn Ala Val Gly Ala Gly Thr
       160  65                  70                  75                  80
       161 Gly Ile Val Ile Asp Pro Asn Gly Val Val Leu Thr Asn Asn His Val
       162              85                  90                  95
       163 Ile Ala Gly Ala Thr Asp Ile Asn Ala Phe Ser Val Gly Ser Gly Gln
       164              100                 105                 110
       165 Thr Tyr Gly Val Asp Val Val Gly Tyr Asp Arg Thr Gln Asp Val Ala
       166          115                 120                 125
       167 Val Leu Gln Leu Arg Gly Ala Gly Gly Leu Pro Ser Ala Ala Ile Gly
       168      130                 135                 140
       169 Gly Gly Val Ala Val Gly Glu Pro Val Val Ala Met Gly Asn Ser Gly
       170 145                 150                 155                 160
       171 Gly Gln Gly Gly Thr Pro Arg Ala Val Pro Gly Arg Val Val Ala Leu
       172              165                 170                 175
       173 Gly Gln Thr Val Gln Ala Ser Asp Ser Leu Thr Gly Ala Glu Glu Thr
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Output Set: N:\CRF3\01152002\I597796B.raw

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180 225             230             235                 240
181 Ile Pro Ile Gly Gln Ala Met Ala Ile Ala Gly Gln Ile Arg Ser Gly
182             245             250             255
183 Gly Gly Ser Pro Thr Val His Ile Gly Pro Thr Ala Phe Leu Gly Leu
184             260             265             270
185 Gly Val Val Asp Asn Asn Gly Asn Gly Ala Arg Val Gln Arg Val Val
186      275             280                 285
187 Gly Ser Ala Pro Ala Ala Ser Leu Gly Ile Ser Thr Gly Asp Val Ile
188      290             295             300
189 Thr Ala Val Asp Gly Ala Pro Ile Asn Ser Ala Thr Ala Met Ala Asp
190 305             310                 315                 320
191 Ala Leu Asn Gly His His Pro Gly Asp Val Ile Ser Val Asn Trp Gln
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202 <213> ORGANISM: Mycobacterium tuberculosis
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220 <212> TYPE: PRT
221 <213> ORGANISM: Mycobacterium tuberculosis
223 <220> FEATURE:
224 <223> OTHER INFORMATION: MTBRa12 C-terminus of MTB32A (Ra35FL)
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230             20             25             30
231 Gly Gly Gly Ser Pro Thr Val His Ile Gly Pro Thr Ala Phe Leu Gly
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234      50                      55                      60
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236  65                      70                      75                      80
237 Ile Thr Ala Val Asp Gly Ala Pro Ile Asn Ser Ala Thr Ala Met Ala
238                      85                      90                      95
239 Asp Ala Leu Asn Gly His His Pro Gly Asp Val Ile Ser Val Asn Trp
240              100                     105                     110
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250 <213> ORGANISM: Mycobacterium tuberculosis
252 <220> FEATURE:
253 <223> OTHER INFORMATION: MTB39 (TbH9) cDNA full-length
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258 tgtggctagc aatgccgctg cggcggcccc gacgactggg gtggtgcccc ccgctgccga 180
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<400> 10
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<213> Mycobacterium tuberculosis

<213> Artificial Sequence

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<213> can only have one response:
Artificial Sequence
or Unknown or
Scientific name
(Genus/species)

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<220>
<223> MTB39 (TbH9) cDNA

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Output Set: **N:\CRF3\01152002\I597796B.raw**

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L:132 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:3
L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
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L:508 M:280 W: Numeric Identifier already exists, Organism not replaced.
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